



SUPPLEMENTAL SEQUENCE LISTING FOR 10817530.txt

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<120> PHYSICAL-CHEMICAL PROPERTY BASED SEQUENCE MOTIFS AND METHODS REGARDING SAME

<130> 265.00400101

<140> 10/817,530  
<141> 2004-04-02

<150> US 60/460,769  
<151> 2003-04-04

<160> 17

<170> PatentIn version 3.3

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<213> ARTIFICIAL SEQUENCE

<220>  
<223> MEMBER OF DNase-I SUPERFAMILY

<400> 1

Pro Asp Ile Leu Cys Leu Gln Glu Thr Lys  
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<220>  
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<400> 2

Leu Tyr Glu Asp Pro Pro Asp Gln Lys Thr Ser Pro Ser Gly Lys Pro  
1 5 10 15

Ala Thr Leu Lys Ile Cys Ser Trp Asn Val Asp Gly Leu Arg Ala Trp  
20 25 30

Ile Lys Lys Lys Gly Leu Asp Trp Val Lys Glu Glu Ala Pro Asp Ile  
35 40 45

Leu Cys Leu Gln Glu Thr Lys Cys Ser Glu Asn Lys Leu Pro Ala Glu  
50 55 60

Leu Gln Glu Leu Pro Gly Leu Ser His Gln Tyr Trp Ser Ala Pro Ser  
65 70 75 80

SUPPLEMENTAL SEQUENCE LISTING FOR 10817530.txt

Asp Lys Glu Gly Tyr Ser Gly Val Gly Leu Leu Ser Arg Gln Cys Pro  
85 90 95

Leu Lys Val Ser Tyr Gly Ile Gly Asp Glu Glu His Asp Gln Glu Gly  
100 105 110

Arg Val Ile Val Ala Glu Phe Asp Ser Phe Val Leu Val Thr Ala Tyr  
115 120 125

Val Pro Asn Ala Gly Arg Gly Leu Val Arg Leu Glu Tyr Arg Gln Arg  
130 135 140

Trp Asp Glu Ala Phe Arg Lys Phe Leu Lys Gly Leu Ala Ser Arg Lys  
145 150 155 160

Pro Leu Val Leu Cys Gly Asp Leu Asn Val Ala His Glu Glu Ile Asp  
165 170 175

Leu Arg Asn Pro Lys Gly Asn Lys Lys Asn Ala Gly Phe Thr Pro Gln  
180 185 190

Glu Arg Gln Gly Phe Gly Glu Leu Leu Gln Ala Val Pro Leu Ala Asp  
195 200 205

Ser Phe Arg His Leu Tyr Pro Asn Thr Pro Tyr Ala Tyr Thr Phe Trp  
210 215 220

Thr Tyr Met Met Asn Ala Arg Ser Lys Asn Val Gly Trp Arg Leu Asp  
225 230 235 240

Tyr Phe Leu Leu Ser His Ser Leu Leu Pro Ala Leu Cys Asp Ser Lys  
245 250 255

Ile Arg Ser Lys Ala Leu Gly Ser Asp His Cys Pro Ile Thr Leu Tyr  
260 265 270

Leu Ala Leu  
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<220>  
<223> MEMBER OF DNase-I SUPERFAMILY

<400> 3

Met Lys Phe Val Ser Phe Asn Ile Asn Gly Leu Arg Ala Arg Pro His  
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10

15

Gln Leu Glu Ala Ile Val Glu Lys His Gln Pro Asp Val Ile Gly Leu  
20                   25                   30

Gln Glu Thr Lys Val His Asp Asp Met Phe Pro Leu Glu Glu Val Ala  
35                   40                   45

Lys Leu Gly Tyr Asn Val Phe Tyr His Gly Gln Lys Gly His Tyr Gly  
50                   55                   60

Val Ala Leu Leu Thr Lys Glu Thr Pro Ile Ala Val Arg Arg Gly Phe  
65                   70                   75                   80

Pro Gly Asp Asp Glu Glu Ala Gln Arg Arg Ile Ile Met Ala Glu Ile  
85                   90                   95

Pro Ser Leu Leu Gly Asn Val Thr Val Ile Asn Gly Tyr Phe Pro Gln  
100               105                   110

Gly Glu Ser Arg Asp His Pro Ile Lys Phe Pro Ala Lys Ala Gln Phe  
115               120                   125

Tyr Gln Asn Leu Gln Asn Tyr Leu Glu Thr Glu Leu Lys Arg Asp Asn  
130               135                   140

Pro Val Leu Ile Met Gly Asp Met Asn Ile Ser Pro Thr Asp Leu Asp  
145               150                   155                   160

Ile Gly Ile Gly Glu Glu Asn Arg Lys Arg Trp Leu Arg Thr Gly Lys  
165               170                   175

Cys Ser Phe Leu Pro Glu Glu Arg Glu Trp Met Asp Arg Leu Met Ser  
180               185                   190

Trp Gly Leu Val Asp Thr Phe Arg His Ala Asn Pro Gln Thr Ala Asp  
195               200                   205

Arg Phe Ser Trp Phe Asp Tyr Arg Ser Lys Gly Phe Asp Asp Asn Arg  
210               215                   220

Gly Leu Arg Ile Asp Leu Leu Ala Ser Gln Pro Leu Ala Glu Cys  
225               230                   235                   240

Cys Val Glu Thr Gly Ile Asp Tyr Glu Ile Arg Ser Met Glu Lys Pro  
245               250                   255

Ser Asp His Ala Pro Val Trp Ala Thr Phe Arg Arg

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Leu Lys Ile Ala Ala Phe Asn Ile Arg Thr Phe Gly Glu Thr Lys Met  
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Ser Asn Ala Thr Leu Ala Ser Tyr Ile Val Arg Ile Val Arg Arg Tyr  
20 25 30

Asp Ile Val Leu Ile Gln Glu Val Arg Asp Ser His Leu Val Ala Val  
35 40 45

Gly Lys Leu Leu Asp Tyr Leu Asn Gln Asp Asp Pro Asn Thr Tyr His  
50 55 60

Tyr Val Val Ser Glu Pro Leu Gly Arg Asn Ser Tyr Lys Glu Arg Tyr  
65 70 75 80

Leu Phe Leu Phe Arg Pro Asn Lys Val Ser Val Leu Asp Thr Tyr Gln  
85 90 95

Tyr Asp Asp Gly Cys Cys Gly Asn Asp Ser Phe Ser Arg Glu Pro Ala  
100 105 110

Val Val Lys Phe Ser Ser His Ser Thr Lys Val Lys Glu Phe Ala Ile  
115 120 125

Val Ala Leu His Ser Ala Pro Ser Asp Ala Val Ala Glu Ile Asn Ser  
130 135 140

Leu Tyr Asp Val Tyr Leu Asp Val Gln Gln Lys Trp His Leu Asn Asp  
145 150 155 160

Val Met Leu Met Gly Asp Phe Asn Ala Asp Cys Ser Tyr Val Thr Ser  
165 170 175

Ser Gln Trp Ser Ser Ile Arg Leu Arg Thr Ser Ser Thr Phe Gln Trp  
180 185 190

Leu Ile Pro Asp Ser Ala Asp Thr Thr Ala Thr Ser Thr Asn Cys Ala  
195 200 205

SUPPLEMENTAL SEQUENCE LISTING FOR 10817530.txt

Tyr Asp Arg Ile Val Val Ala Gly Ser Leu Leu Gln Ser Ser Val Val  
210 215 220

Pro Gly Ser Ala Ala Pro Phe Asp Phe Gln Ala Ala Tyr Gly Leu Ser  
225 230 235 240

Asn Glu Met Ala Leu Ala Ile Ser Asp His Tyr Pro Val Glu Val Thr  
245 250 255

Leu Thr

<210> 5  
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<213> ARTIFICIAL SEQUENCE

<220>  
<223> MEMBER OF DNase-I SUPERFAMILY

<400> 5

Tyr Asp Pro Ile His Glu Tyr Val Asn His Glu Leu Arg Lys Arg Glu  
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Asn Glu Phe Ser Glu His Lys Asn Val Lys Ile Phe Val Ala Ser Tyr  
20 25 30

Asn Leu Asn Gly Cys Ser Ala Thr Thr Lys Leu Glu Asn Trp Leu Phe  
35 40 45

Pro Glu Asn Thr Pro Leu Ala Asp Ile Tyr Val Val Gly Phe Gln Glu  
50 55 60

Ile Val Gln Leu Thr Ser Ala Asp Pro Ala Lys Arg Arg Glu Trp Glu  
65 70 75 80

Ser Cys Val Lys Arg Leu Leu Asn Gly Lys Cys Thr Ser Gly Pro Gly  
85 90 95

Tyr Val Gln Leu Arg Ser Gly Gln Leu Val Gly Thr Ala Leu Met Ile  
100 105 110

Phe Cys Lys Glu Ser Cys Leu Pro Ser Ile Lys Asn Val Glu Gly Thr  
115 120 125

Val Lys Lys Thr Gly Leu Gly Asn Lys Gly Ala Val Ala Ile Arg Phe  
130 135 140

Asp Tyr Glu Asp Thr Gly Leu Cys Phe Ile Thr Ser His Leu Ala Ala

145

SUPPLEMENTAL SEQUENCE LISTING FOR 10817530.txt  
150 155 160

Gly Tyr Thr Asn Tyr Asp Glu Arg Asp His Asp Tyr Arg Thr Ile Ala  
165 170 175

Ser Gly Leu Arg Phe Arg Arg Gly Arg Ser Ile Phe Asn His Asp Tyr  
180 185 190

Val Val Trp Phe Gly Asp Phe Asn Tyr Arg Ile Ser Leu Thr Tyr Glu  
195 200 205

Glu Val Val Pro Cys Ile Ala Gln Gly Lys Leu Ser Tyr Leu Phe Glu  
210 215 220

Tyr Asp Gln Leu Asn Lys Gln Met Leu Thr Gly Lys Val Phe Pro Phe  
225 230 235 240

Phe Ser Glu Leu Pro Ile Thr Phe Pro Pro Thr Tyr Lys Phe Asp Ile  
245 250 255

Gly Thr Asp Ile Tyr Asp Thr Ser Asp Lys His Arg Val Pro Ala Trp  
260 265 270

Thr Asp Arg Ile Leu Tyr Arg Gly Glu Leu Val Pro His Ser Tyr Gln  
275 280 285

Ser Val Pro Leu Tyr Tyr Ser Asp His Arg Pro Ile Tyr Ala Thr Tyr  
290 295 300

Glu Ala Asn Ile Val Lys Val Asp Arg Glu Lys Lys Lys Ile Leu Phe  
305 310 315 320

Glu Glu Leu Tyr Asn Gln Arg Lys Gln Glu Val Arg Asp Ala Ser Gln  
325 330 335

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<212> PRT

<213> HOMO SAPIENS

<400> 6

Leu Lys Ile Cys Ser Trp Asn Val Asp Gly Leu Arg Ala  
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<210> 7

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SUPPLEMENTAL SEQUENCE LISTING FOR 10817530.txt

Pro Asp Ile Leu Cys Leu Gln Glu Thr Lys  
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Lys Glu Gly Tyr Ser Gly Val Gly Leu Leu Ser Arg Gln Cys Pro  
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Gly Ile Gly Asp Glu Glu His Asp Gln Glu Gly Arg Val Ile Val Ala  
1 5 10 15

Glu Phe Asp Ser Phe Val Leu  
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Tyr Val Pro Asn Ala  
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Arg Leu Glu Tyr Arg Gln Arg Trp  
1 5

<210> 12  
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Pro Leu Val Leu Cys Gly Asp Leu Asn Val Ala His  
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SUPPLEMENTAL SEQUENCE LISTING FOR 10817530.txt

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Gly Phe Thr Pro Gln Glu Arg Gln Gly Phe Gly Glu Leu  
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Val Pro Leu Ala Asp Ser Phe Arg  
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<400> 15

Tyr Thr Phe Trp Thr Tyr Met  
1 5

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Arg Ser Lys Asn Val Gly Trp Arg Leu Asp Tyr Phe Leu Leu Ser His  
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Ser Leu

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Gly Ser Asp His Cys Pro Ile  
1 5